

SEQUENCE PROTOCOL

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tctgaggacg ccgactctgc agatgcagac aacgcgagca atgtaatcaa tgagaatgag 180

gactcctcgg aaggtgctaa ccagccttca aacgagtcac cctctacgga agccaaatcc 240

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40 gtg ggt tac gaa act cct tcc cca att cag gca caa acc atc cca atc 339
 Val Gly Tyr Glu Thr Pro Ser Pro Ile Gln Ala Gln Thr Ile Pro Ile
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50 aag act gca gct ttc gcg ctg cca atc ctt gcc cgt att gac aag tcc 435
 Lys Thr Ala Ala Phe Ala Leu Pro Ile Leu Ala Arg Ile Asp Lys Ser
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55 gtg cgc agc cca cag gca ctt gtg ctt gcc cct acc cgt gag cag gca 483
 Val Arg Ser Pro Gln Ala Leu Val Leu Ala Pro Thr Arg Glu Gln Ala
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55 ctt cag gtt gct gac tcc ttc caa tcc ttc gct gac cac gtc ggt ggc 531
 Leu Gln Val Ala Asp Ser Phe Gln Ser Phe Ala Asp His Val Gly Gly
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5	tct ggc ctg cgt cgt ggc gct cac atc gtc gtg ggt acc cca ggc cga	627
	Ser Gly Leu Arg Arg Gly Ala His Ile Val Val Gly Thr Pro Gly Arg	
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10	atc atc gat cac ctc gaa aag ggc tcc ctg gat atc tcc gga ctg cgc	675
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15	ttc ctc gtg ctc gat gaa gca gac gag atg ctg aac atg ggc ttc cag	723
	Phe Leu Val Leu Asp Glu Ala Asp Glu Met Leu Asn Met Gly Phe Gln	
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	Glu Asp Val Glu Arg Ile Leu Glu Asp Thr Pro Asp Glu Lys Gln Val	
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20	aag gaa tac tcc cag gcc aac gac gtt cct cta gag gac atc gca gcg	1491
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55	atc acc atc gca gcc gac cac acc ctg gtt gaa ctg cca aag gat ctc	1827
	Ile Thr Ile Ala Ala Asp His Thr Leu Val Glu Leu Pro Lys Asp Leu	
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	525 530 535	
65	ctc atc aac ata gaa cgc gac tcc ggt gga cgc cca cca cgc cgc ttc	1923
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10 ggc ttc cgc ggt gga cgt gac cgt gat gat cgt ggc gga cgc ggt gga 2067
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 590 595 600

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20 cgt ggc gga cgc gac taagagttcg ttttagcttc agctcagggtt ttcgcctgag 2170
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50 Ser Phe Gln Ser Phe Ala Asp His Val Gly Gly Leu Asn Val Leu Pro
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000557 BT / AL1